WEST Search History for Application 09539032

Creation Date: 2008121918:41

Query	DB	Op.	Plur.	Thes.	Date
dash.in. and peptide.clm.	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
brahmachari.in.	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(brahmachari.in.) and peptide.clm.	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
sequence adj alignment and overlap	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap) and (protein or peptide) and (pathogen or bacteria or virus or fungus)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus)) and sequence same (compare or comparison)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison)) and (computer or computerized or automate or automated)	PGPB, USPT, EPAB, JPAB,	OR	YES		12-19-2008

	DWPI, TDBD				
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated)) and fdrug adj target	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated)) and drug adj target	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated) and drug adj target) and conserved adj (protein or peptide)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated) and drug adj target and conserved adj (protein or peptide)) and (bacillus adj subtilis or mycoplasma adj pneumoniae or helicobacter adj pyloir)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated) and drug adj target and conserved adj (protein or peptide)) and helicobacter adj pylori	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated) and drug adj target and conserved adj (protein or peptide)) and bacillus adj subtilis	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated) and drug adj target and conserved adj (protein or peptide)) and hemophilus adj influenzae	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008

(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated) and drug adj target and conserved adj (protein or peptide)) and escherichia coli	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(sequence adj alignment and overlap and (protein or peptide) and (pathogen or bacteria or virus or fungus) and sequence same (compare or comparison) and (computer or computerized or automate or automated) and drug adj target and conserved adj (protein or peptide)) and escherichia adj coli	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
motif or pattern	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
same alignmentL17	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern) same alignment	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment) and (microorganism or pathogen or microb$)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$)) and (invariant or conserved or common)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common)) and (peptide or protein or polypeptide)	PGPB, USPT, EPAB, JPAB,	OR	YES		12-19-2008

	DWPI, TDBD				
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common) and (peptide or protein or polypeptide)) and host	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common) and (peptide or protein or polypeptide) and host) and database	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common) and (peptide or protein or polypeptide) and host and database) and drug adj target	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common) and (peptide or protein or polypeptide) and host and database and drug adj target) and (computer or computerized or automate or automated)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common) and (peptide or protein or polypeptide) and host and database and drug adj target and (computer or computerized or automate or automated)) and (matched or matching)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common) and (peptide or protein or polypeptide) and host and database and drug adj target and (computer or computerized or automate or automated) and (matched or matching)) and protein adj identification adj number	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(motif or pattern same alignment and (microorganism or pathogen or microb$) and (invariant or conserved or common) and (peptide or protein or polypeptide) and host and database and drug adj target and (computer or computerized or automate or automated) and (matched or matching)) and annotation	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
comparative adh genomics		OR	YES		12-19-2008

	PGPB, USPT, EPAB, JPAB, DWPI, TDBD				
comparative adj genomics	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(comparative adj genomics) and conserved adj peptide adj motif	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
drug adj screening	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(drug adj screening) and (pathogen or bacteria)	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(drug adj screening and (pathogen or bacteria)) and protein adj alignment	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008
(drug adj screening and (pathogen or bacteria) and protein adj alignment) and bacterial adj genome	PGPB, USPT, EPAB, JPAB, DWPI, TDBD	OR	YES		12-19-2008